Application No. 10/516,741 Docket No.: 4528-0109PUS2

Amendment dated February 5, 2008 After Final Office Action of November 5, 2007

AMENDMENTS TO THE CLAIMS

 (Currently Amended) A method for identifying heterologous DNA and the mRNA transcribed therefrom, which causes, on its expression, an electrophysiological change in a cell comprising the steps of:

- (i) providing a substrate for making the electrophysiological measurements upon which at least one cell can be arranged;
- (ii) providing a plurality of cells, each cell comprising a different heterologous DNA sequence, said DNA sequence being a member of a DNA library; wherein each cell expresses the heterologous DNA sequence it comprises:
- (iii) arranging the plurality of cells provided in step (ii) on the substrate to permit simultaneous detection and/or measurement of a change (in comparison to a control cell) in the electrophysiology of each cell, said change being a result of expression of the heterologous DNA sequence, and
- (iv) identifying at least one cell of interest, which shows a change in its electrophysiology electrophysiology as measured in step (iii), characterized in that, the method comprises the further steps of:

isolating the cell of interest, and/or genetic material therefrom; and isolating mRNA from the cell of interest showing a change in its electrophysiology as measured in step (iii), wherein the mRNA is isolated without removing the cell from the substrate.

- (Previously Presented) The method as claimed in Claim 1, wherein the method further comprises the step of sequencing the genetic material.
- 3. (Withdrawn) The method as claimed in Claim 2, wherein the method further comprises the step of storing or recording the sequence information on an information carrier.

4. - 6. (Cancelled)

Application No. 10/516,741 Docket No.: 4528-0109PUS2 Amendment dated February 5, 2008

After Final Office Action of November 5, 2007

7. (Previously Presented) The method as claimed in Claim 1, wherein the DNA library is

a cDNA library.

8. (Previously Presented) The method as claimed in Claim 1, wherein the change in the

electrophysiology of the cell is detected and/or measured by patch clamping.

9. (Previously Presented) The method as claimed in Claim 1, wherein the cell is treated

with a test agent before step (iii).

10. (Previously Presented) The method as claimed in Claim 9, wherein the test agent is

selected from at least one of the following: small organic molecules, small peptides,

neurotransmitters, hormones and cytokines.

11. (Previously Presented) The method as claimed in Claim 1, wherein the cell is an

animal cell.

12. (Previously Presented) The method as claimed in Claim 1, wherein the animal cell is

selected from: Human Embryonic Kidney 293 (HEK293), Chinese Hamster Ovary (CHO), COS,

MDCK, NG108, NIH3T3 or T84.

13. (Previously Presented) The method as claimed in Claim 1, wherein the cells are

arranged at spaced-apart locations on the substrate.

14. (Withdrawn) The method as claimed in Claim 3, wherein said information carrier is a

3

computer disk.

GMM/MTC/